

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

10/517,220

Source:

PCT

Date Processed by STIC:

5-19-05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05

Raw Sequence Listing Error Summary

ERROR DETECTED
SUGGESTED CORRECTION
SERIAL NUMBER:

10/517,220

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1	____ Wrapped Nucleic- Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2	____ Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3	____ Misaligned Amino Numbering	The <u>numbering under each 5th amino acid</u> is misaligned. <u>Do not use tab codes between numbers:</u> use <u>space characters</u> , instead.
4	____ Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5	____ Variable Length	Sequence(s) ____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6	____ PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7	____ Skipped Sequences (OLD RULES)	Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8	____ Skipped Sequences (NEW RULES)	Sequence(s) ____ missing. If intentional , please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9	____ Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
10	____ Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
11	____ Use of <220>	Sequence(s) ____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
12	____ PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13	____ Misuse of n/Xaa	"n" can only represent a single <u>nucleotide</u> ; "Xaa" can only represent a single <u>amino acid</u>



PCT

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/517,220

DATE: 07/19/2005
TIME: 08:58:58

Input Set : E:\24615-20244.00 - seq list.txt
Output Set: N:\CRF4\07192005\J517220.raw

3 <110> APPLICANT: Edens, Lupo
4 Lopez, Michel
6 <120> TITLE OF INVENTION: Improved method for the prevention or reduction of haze in
7 beverages
9 <130> FILE REFERENCE: 246152024400
11 <140> CURRENT APPLICATION NUMBER: US 10/517,220
C--> 12 <141> CURRENT FILING DATE: 2004-12-07
14 <150> PRIOR APPLICATION NUMBER: PCT/NL03/00352
15 <151> PRIOR FILING DATE: 2003-05-14
17 <150> PRIOR APPLICATION NUMBER: NL 02100681.2
18 <151> PRIOR FILING DATE: 2002-06-07
20 <160> NUMBER OF SEQ ID NOS: 7
22 <170> SOFTWARE: PatentIn Ver. 2.1

Does Not Comply
Corrected Diskette Needed
(Pg. 1-S) 2

ERRORED SEQUENCES

260 <210> SEQ ID NO: 4
261 <211> LENGTH: 526
262 <212> TYPE: PRT
263 <213> ORGANISM: Aspergillus niger
265 <400> SEQUENCE: 4
266 Met Arg Ser Phe Ser Val Val Ala Ala Ala Ser Leu Ala Leu Ser Trp
E--> 267 1 5 10 15
268 Ala Ser Leu Ala Gln Ala Ala Arg Pro Arg Leu Val Pro Lys Pro Ile
E--> 269 20 25 30
270 Ser Arg Pro Ala Ser Ser Lys Ser Ala Ala Thr Thr Gly Glu Ala Tyr
E--> 271 35 40 45
272 Phe Glu Gln Leu Leu Asp His His Asn Pro Glu Lys Gly Thr Phe Ser
E--> 273 50 55 60
274 Gln Arg Tyr Trp Trp Ser Thr Glu Tyr Trp Gly Gly Pro Gly Ser Pro
E--> 275 65 70 75 80
276 Val Val Leu Phe Asn Pro Gly Glu Val Ser Ala Asp Gly Tyr Glu Gly
E--> 277 85 90 95
278 Tyr Leu Thr Asn Asp Thr Leu Thr Gly Val Tyr Ala Gln Glu Ile Gln
E--> 279 100 105 110
280 Gly Ala Val Ile Leu Ile Glu His Arg Tyr Trp Gly Asp Ser Ser Pro
E--> 281 115 120 125
282 Tyr Glu Val Leu Asn Ala Glu Thr Leu Gln Tyr Leu Thr Leu Asp Gln
E--> 283 130 135 140
284 Ser Ile Leu Asp Met Thr Tyr Phe Ala Glu Thr Val Lys Leu Gln Phe
E--> 285 145 150 155 160
286 Asp Asn Ser Ser Arg Ser Asn Ala Gln Asn Ala Pro Trp Val Met Val

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See item
3 on
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RAW SEQUENCE LISTING
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Input Set : E:\24615-20244.00 - seq list.txt
Output Set: N:\CRF4\07192005\J517220.raw

E--> 287 165 170 175
288 Gly Gly Ser Tyr Ser Gly Ala Leu Thr Ala Trp Thr Glu Ser Ile Ala
E--> 289 180 185 190
290 Pro Gly Thr Phe Trp Ala Tyr His Ala Thr Ser Ala Pro Val Glu Ala
E--> 291 195 200 205
292 Ile Tyr Asp Phe Trp Gln Tyr Phe Tyr Pro Ile Gln Gln Gly Met Ala
E--> 293 210 215 220
294 Gln Asn Cys Ser Lys Asp Val Ser Leu Val Ala Glu Tyr Val Asp Lys
E--> 295 225 230 235 240
296 Ile Gly Lys Asn Gly Thr Ala Lys Glu Gln Gln Glu Leu Lys Glu Leu
E--> 297 245 250 255
298 Phe Gly Leu Gly Ala Val Glu His Tyr Asp Asp Phe Ala Ala Val Leu
E--> 299 260 265 270
300 Pro Asn Gly Pro Tyr Leu Trp Gln Asp Asn Asp Phe Val Thr Gly Tyr
E--> 301 275 280 285
302 Ser Ser Phe Phe Gln Phe Cys Asp Ala Val Glu Gly Val Glu Ala Gly
E--> 303 290 295 300
304 Ala Ala Val Thr Pro Gly Pro Glu Gly Val Gly Leu Glu Lys Ala Leu
E--> 305 305 310 315 320
306 Ala Asn Tyr Ala Asn Trp Phe Asn Ser Thr Ile Leu Pro Asn Tyr Cys
E--> 307 325 330 335
308 Ala Ser Tyr Gly Tyr Trp Thr Asp Glu Trp Ser Val Ala Cys Phe Asp
E--> 309 340 345 350
310 Ser Tyr Asn Ala Ser Ser Pro Ile Phe Thr Asp Thr Ser Val Gly Asn
E--> 311 355 360 365
312 Pro Val Asp Arg Gln Trp Glu Trp Phe Leu Cys Asn Glu Pro Phe Phe
E--> 313 370 375 380
314 Trp Trp Gln Asp Gly Ala Pro Glu Gly Thr Ser Thr Ile Val Pro Arg
E--> 315 385 390 395 400
316 Leu Val Ser Ala Ser Tyr Trp Gln Arg Gln Cys Pro Leu Tyr Phe Pro
E--> 317 405 410 415
318 Glu Val Asn Gly Tyr Thr Tyr Gly Ser Ala Lys Gly Lys Asn Ser Ala
E--> 319 420 425 430
320 Thr Val Asn Ser Trp Thr Gly Gly Trp Asp Met Thr Arg Asn Thr Thr
E--> 321 435 440 445
322 Arg Leu Ile Trp Thr Asn Gly Gln Tyr Asp Pro Trp Arg Asp Ser Gly
E--> 323 450 455 460
324 Val Ser Ser Thr Phe Arg Pro Gly Gly Pro Leu Val Ser Thr Ala Asn
E--> 325 465 470 475 480
326 Glu Pro Val Gln Ile Ile Pro Gly Gly Phe His Cys Ser Asp Leu Tyr
E--> 327 485 490 495
328 Met Glu Asp Tyr Tyr Ala Asn Glu Gly Val Arg Lys Val Val Asp Asn
E--> 329 500 505 510
330 Glu Val Lys Gln Ile Lys Glu Trp Val Glu Glu Tyr Tyr Ala
E--> 331 515 520 525
333 <210> SEQ ID NO: 5
334 <211> LENGTH: 526
335 <212> TYPE: PRT
336 <213> ORGANISM: Aspergillus niger

CPIS
See item
3 on
error
Summary
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/517,220

DATE: 07/19/2005
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Input Set : E:\24615-20244.00 - seq list.txt
Output Set: N:\CRF4\07192005\J517220.raw

338 <400> SEQUENCE: 5
 339 Met Arg Ala Phe Ser Ala Val Ala Ala Ala Leu Ala Leu Ser Trp
 E--> 340 1 5 10 15
 341 Ala Ser Leu Ala Gln Ala Ala Arg Pro Arg Leu Val Pro Lys Pro Val
 E--> 342 20 25 30
 343 Ser Arg Pro Ala Ser Ser Lys Ser Ala Ala Thr Thr Gly Glu Ala Tyr
 E--> 344 35 40 45
 345 Phe Glu Gln Leu Leu Asp His His Asn Pro Glu Lys Gly Thr Phe Ser
 E--> 346 50 55 60
 347 Gln Arg Tyr Trp Trp Ser Thr Glu Tyr Trp Gly Gly Pro Gly Ser Pro
 E--> 348 65 70 75 80
 349 Val Val Leu Phe Thr Pro Gly Glu Val Ser Ala Asp Gly Tyr Glu Gly
 E--> 350 85 90 95
 351 Tyr Leu Thr Asn Gly Thr Leu Thr Gly Val Tyr Ala Gln Glu Ile Gln
 E--> 352 100 105 110
 353 Gly Ala Val Ile Leu Ile Glu His Arg Tyr Trp Gly Asp Ser Ser Pro
 E--> 354 115 120 125
 355 Tyr Glu Val Leu Asn Ala Glu Thr Leu Gln Tyr Leu Thr Leu Asp Gln
 E--> 356 130 135 140
 357 Ala Ile Leu Asp Met Thr Tyr Phe Ala Glu Thr Val Lys Leu Gln Phe
 E--> 358 145 150 155 160
 359 Asp Asn Ser Thr Arg Ser Asn Ala Gln Asn Ala Pro Trp Val Met Val
 E--> 360 165 170 175
 361 Gly Gly Ser Tyr Ser Gly Ala Leu Thr Ala Trp Thr Glu Ser Val Ala
 E--> 362 180 185 190
 363 Pro Gly Thr Phe Trp Ala Tyr His Ala Thr Ser Ala Pro Val Glu Ala
 E--> 364 195 200 205
 365 Ile Tyr Asp Tyr Trp Gln Tyr Phe Tyr Pro Ile Gln Gln Gly Met Ala
 E--> 366 210 215 220
 367 Gln Asn Cys Ser Lys Asp Val Ser Leu Val Ala Glu Tyr Val Asp Lys
 E--> 368 225 230 235 240
 369 Ile Gly Lys Asn Gly Thr Ala Lys Glu Gln Gln Ala Leu Lys Glu Leu
 E--> 370 245 250 255
 371 Phe Gly Leu Gly Ala Val Glu His Phe Asp Asp Phe Ala Ala Val Leu
 E--> 372 260 265 270
 373 Pro Asn Gly Pro Tyr Leu Trp Gln Asp Asn Asp Phe Ala Thr Gly Tyr
 E--> 374 275 280 285
 375 Ser Ser Phe Phe Gln Phe Cys Asp Ala Val Glu Gly Val Glu Ala Gly
 E--> 376 290 295 300
 377 Ala Ala Val Thr Pro Gly Pro Glu Gly Val Gly Leu Glu Lys Ala Leu
 E--> 378 305 310 315 320
 379 Ala Asn Tyr Ala Asn Trp Phe Asn Ser Thr Ile Leu Pro Asp Tyr Cys
 E--> 380 325 330 335
 381 Ala Ser Tyr Gly Tyr Trp Thr Asp Glu Trp Ser Val Ala Cys Phe Asp
 E--> 382 340 345 350
 383 Ser Tyr Asn Ala Ser Ser Pro Ile Tyr Thr Asp Thr Ser Val Gly Asn
 E--> 384 355 360 365
 385 Ala Val Asp Arg Gln Trp Glu Trp Phe Leu Cys Asn Glu Pro Phe Phe
 E--> 386 370 375 380

-same
error

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Input Set : E:\24615-20244.00 - seq list.txt
Output Set: N:\CRF4\07192005\J517220.raw

387 Tyr Trp Gln Asp Gly Ala Pro Glu Gly Thr Ser Thr Ile Val Pro Arg
E--> 388 385 390 395 400
389 Leu Val Ser Ala Ser Tyr Trp Gln Arg Gln Cys Pro Leu Tyr Phe Pro
E--> 390 405 410 415
391 Glu Thr Asn Gly Tyr Thr Tyr Gly Ser Ala Lys Gly Lys Asn Ala Ala
E--> 392 420 425 430
393 Thr Val Asn Ser Trp Thr Gly Gly Trp Asp Met Thr Arg Asn Thr Thr
E--> 394 435 440 445
395 Arg Leu Ile Trp Thr Asn Gly Gln Tyr Asp Pro Trp Arg Asp Ser Gly
E--> 396 450 455 460
397 Val Ser Ser Thr Phe Arg Pro Gly Gly Pro Leu Ala Ser Thr Ala Asn
E--> 398 465 470 475 480
399 Glu Pro Val Gln Ile Ile Pro Gly Gly Phe His Cys Ser Asp Leu Tyr
E--> 400 485 490 495
401 Met Ala Asp Tyr Tyr Ala Asn Glu Gly Val Lys Lys Val Val Asp Asn
E--> 402 500 505 510
403 Glu Val Lys Gln Ile Lys Glu Trp Val Glu Glu Tyr Tyr Ala
E--> 404 515 520 525
464 <210> SEQ ID NO: 7
465 <211> LENGTH: 516
466 <212> TYPE: PRT
467 <213> ORGANISM: Aspergillus niger
469 <400> SEQUENCE: 7
470 Met Arg Ser Phe Ser Val Val Ala Ala Ser Leu Ala Leu Ser Trp
E--> 471 1 5 10 15
472 Ala Ser Leu Ala Gln Ala Ala Arg Pro Arg Leu Val Pro Lys Pro Ile
E--> 473 20 25 30
474 Ser Arg Pro Ala Ser Ser Lys Ser Ala Ala Thr Thr Gly Glu Ala Tyr
E--> 475 35 40 45
476 Phe Glu Gln Leu Leu Asp His His Asn Pro Glu Lys Gly Thr Phe Ser
E--> 477 50 55 60
478 Gln Arg Tyr Trp Trp Ser Thr Glu Tyr Trp Gly Gly Pro Gly Ser Pro
E--> 479 65 70 75 80
480 Val Val Leu Phe Asn Pro Gly Glu Val Ser Ala Asp Gly Tyr Glu Gly
E--> 481 85 90 95
482 Tyr Leu Thr Asn Asp Thr Leu Thr Gly Val Tyr Ala Gln Glu Ile Gln
E--> 483 100 105 110
484 Gly Ala Val Ile Leu Ile Glu His Arg Tyr Trp Gly Asp Ser Ser Pro
E--> 485 115 120 125
486 Tyr Glu Val Leu Asn Ala Glu Thr Leu Gln Tyr Leu Thr Leu Asp Gln
E--> 487 130 135 140
488 Ser Ile Leu Asp Met Thr Tyr Phe Ala Glu Thr Val Lys Leu Gln Phe
E--> 489 145 150 155 160
490 Asp Asn Ser Ser Arg Ser Asn Ala Gln Asn Ala Pro Trp Val Met Val
E--> 491 165 170 175
492 Gly Gly Ser Tyr Ser Gly Ala Leu Thr Ala Trp Thr Glu Ser Ile Ala
E--> 493 180 185 190
494 Pro Gly Thr Phe Trp Ala Tyr His Ala Thr Ser Ala Pro Val Glu Ala
E--> 495 195 200 205

*/SAME
errors*

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PATENT APPLICATION: US/10/517,220

DATE: 07/19/2005
TIME: 08:58:58

Input Set : E:\24615-20244.00 - seq list.txt
Output Set: N:\CRF4\07192005\J517220.raw

496 Ile Tyr Asp Phe Gln Gly Met Ala Gln Asn Cys Ser Lys Asp Val Ser
E--> 497 210 215 220
498 Leu Val Ala Glu Tyr Val Asp Lys Ile Gly Lys Asn Gly Thr Ala Lys
E--> 499 225 230 235 240
500 Glu Gln Gln Glu Leu Lys Glu Leu Phe Gly Leu Gly Ala Val Glu His
E--> 501 245 250 255
502 Tyr Asp Asp Phe Ala Ala Val Leu Pro Asn Gly Pro Tyr Leu Trp Gln
E--> 503 260 265 270
504 Asp Asn Asp Phe Val Thr Gly Tyr Ser Ser Phe Phe Gln Phe Cys Asp
E--> 505 275 280 285
506 Ala Val Glu Gly Val Glu Ala Gly Ala Ala Val Thr Pro Gly Pro Glu
E--> 507 290 295 300
508 Gly Val Gly Leu Glu Lys Ala Leu Ala Asn Tyr Ala Asn Trp Phe Asn
E--> 509 305 310 315 320
510 Ser Thr Ile Leu Pro Asn Tyr Cys Ala Ser Tyr Gly Tyr Trp Thr Asp
E--> 511 325 330 335
512 Glu Trp Ser Val Ala Cys Phe Asp Ser Tyr Asn Ala Ser Ser Pro Ile
E--> 513 340 345 350
514 Phe Thr Asp Thr Ser Val Gly Asn Pro Val Asp Arg Gln Trp Glu Trp
E--> 515 355 360 365
516 Phe Leu Cys Asn Glu Pro Phe Trp Trp Gln Asp Gly Ala Pro Glu
E--> 517 370 375 380
518 Gly Thr Ser Thr Ile Val Pro Arg Leu Val Ser Ala Ser Tyr Trp Gln
E--> 519 385 390 395 400
520 Arg Gln Cys Pro Leu Tyr Phe Pro Glu Val Asn Gly Tyr Thr Tyr Gly
E--> 521 405 410 415
522 Ser Ala Lys Gly Lys Asn Ser Ala Thr Val Asn Ser Trp Thr Gly Gly
E--> 523 420 425 430
524 Trp Asp Met Thr Arg Asn Thr Thr Arg Leu Ile Trp Thr Asn Gly Gln
E--> 525 435 440 445
526 Tyr Asp Pro Trp Arg Asp Ser Gly Val Ser Ser Thr Phe Arg Pro Gly
E--> 527 450 455 460
528 Gly Pro Leu Val Ser Thr Ala Asn Glu Pro Val Gln Ile Ile Pro Gly
E--> 529 465 470 475 480
530 Gly Phe His Cys Ser Asp Leu Tyr Met Glu Asp Tyr Tyr Ala Asn Glu
E--> 531 485 490 495
532 Gly Val Arg Lys Val Val Asp Asn Glu Val Lys Gln Ile Lys Glu Tyr
E--> 533 500 505 510
534 Gly Tyr Gly Cys
E--> 535 515

- Same
version

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/517,220

DATE: 07/19/2005
TIME: 08:58:59

Input Set : E:\24615-20244.00 - seq list.txt
Output Set: N:\CRF4\07192005\J517220.raw

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:267 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:4
M:332 Repeated in SeqNo=4
L:340 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:5
M:332 Repeated in SeqNo=5
L:471 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
M:332 Repeated in SeqNo=7